

Serial Number: 10/020,478**ENTERED**Entered by: 2Verified by: 2

(STIC stat)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: #2
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEO ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEO ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☒ Deleted extra invalid headings used by an applicant, specifically: 2207 in seq. 3
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/lastname at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/195

OIPE

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/10/020,478

DATE: 01/07/2002  
 TIME: 20:32:06

Input Set : A:\pto.txt  
 Output Set: N:\CRF3\01072002\J020478.raw

# 2.

3 <110> APPLICANT: C. Frank Bennett  
 4 Kenneth Dobie  
 6 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF B-CELL ASSOCIATED PROTEIN EXPRESSION  
 8 <130> FILE REFERENCE: RTS-0303  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/020,478  
 C--> 10 <141> CURRENT FILING DATE: 2001-12-13  
 10 <160> NUMBER OF SEQ ID NOS: 88  
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 46 <222> LOCATION: (186)...(1085)  
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 51 ctttctttcg ccagccttac gggcccgaac cctcgtgtga aggggtcagt acctaagccg 120  
 53 gagcggggta gaggcgggcc ggcacccctc tctgacctcc agtgccgccg gcctcaagat 180  
 55 cagac atg gcc cag aac ttg aag gac ttg gcg gga cgg ctg ccc gcc ggg 230  
 56 Met Ala Gln Asn Leu Lys Asp Leu Ala Gly Arg Leu Pro Ala Gly  
 57 1 5 10 15  
 59 ccc cgg ggc atg ggc acg gcc ctg aag ctg ttg ctg ggg gcc ggc gcc 278  
 60 Pro Arg Gly Met Gly Thr Ala Leu Lys Leu Leu Gly Ala Gly Ala  
 61 20 25 30  
 63 gtg gcc tac ggt gtg cgc gaa tct gtg ttc acc gtg gaa ggc ggg cac 326  
 64 Val Ala Tyr Gly Val Arg Glu Ser Val Phe Thr Val Glu Gly Gly His  
 65 35 40 45  
 67 aga gcc atc ttc ttc aat cgg atc ggt gga gtg cag cag gac act atc 374  
 68 Arg Ala Ile Phe Phe Asn Arg Ile Gly Gly Val Gln Gln Asp Thr Ile  
 69 50 55 60  
 71 ctg gcc gag ggc ctt cac ttc agg atc cct tgg ttc cag tac ccc att 422

RAW SEQUENCE LISTING  
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75 atc tat gac att cgg gcc aga cct cga aaa atc tcc tcc cct aca ggc      470
76 Ile Tyr Asp Ile Arg Ala Arg Pro Arg Lys Ile Ser Ser Pro Thr Gly
77 80      85              90
79 tcc aaa gac cta cag atg gtg aat atc tcc ctg cga gtg ttg tct cga      518
80 Ser Lys Asp Leu Gln Met Val Asn Ile Ser Leu Arg Val Leu Ser Arg
81      100              105
83 ccc aat gct cag gag ctt cct agc atg tac cag cgc cta ggg ctg gac      566
84 Pro Asn Ala Gln Glu Leu Pro Ser Met Tyr Gln Arg Leu Gly Leu Asp
85      115              120
87 tac gag gaa cga gtg ttg ccg tcc att gtc aac gag gtg ctc aag agt      614
88 Tyr Glu Glu Arg Val Leu Pro Ser Ile Val Asn Glu Val Leu Lys Ser
89      130              135
91 gtg gtg gcc aag ttc aat gcc tca cag ctg atc acc cag cgg gcc cag      662
92 Val Val Ala Lys Phe Asn Ala Ser Gln Leu Ile Thr Gln Arg Ala Gln
93      145              150
95 gta tcc ctg ttg atc cgc cgg gag ctg aca gag agg gcc aag gac ttc      710
96 Val Ser Leu Leu Ile Arg Arg Glu Leu Thr Glu Arg Ala Lys Asp Phe
97 160      165              170
99 agc ctc atc ctg gat gat gtg gcc atc aca gag ctg agc ttt agc cga      758
100 Ser Leu Ile Leu Asp Asp Val Ala Ile Thr Glu Leu Ser Phe Ser Arg
101      180              185
103 gag tac aca gct gct gta gaa gcc aaa caa gtg gcc cag cag gag gcc      806
104 Glu Tyr Thr Ala Ala Val Glu Ala Lys Gln Val Ala Gln Gln Glu Ala
105      195              200
107 cag cgg gcc caa ttc ttg gta gaa aaa gca aag cag gaa cag cgg cag      854
108 Gln Arg Ala Gln Phe Leu Val Glu Lys Ala Lys Gln Glu Gln Arg Gln
109      210              215
111 aaa att gtg cag gcc gag ggt gag gcc gag gct gcc aag atg ctt gga      902
112 Lys Ile Val Gln Ala Glu Gly Glu Ala Glu Ala Ala Lys Met Leu Gly
113      225              230
115 gaa gca ctg agc aag aac cct ggc tac atc aaa ctt cgc aag att cga      950
116 Glu Ala Leu Ser Lys Asn Pro Gly Tyr Ile Lys Leu Arg Lys Ile Arg
117 240      245              250
119 gca gcc cag aat atc tcc aag acg atc gcc aca tca cag aat cgt atc      998
120 Ala Ala Gln Asn Ile Ser Lys Thr Ile Ala Thr Ser Gln Asn Arg Ile
121      260              265
123 tat ctc aca gct gac aac ctt gtg ctg aac cta cag gat gaa agt ttc      1046
124 Tyr Leu Thr Ala Asp Asn Leu Val Leu Asn Leu Gln Asp Glu Ser Phe
125      275              280
127 acc agg gga agt gac agc ctc atc aag ggt aag aaa tga gcctagtcac      1095
128 Thr Arg Gly Ser Asp Ser Leu Ile Lys Gly Lys Lys
129      290              295
131 caagaactcc acccccagag gaagtggatc tgcttctcca gtttttgagg agccagccag      1155
133 ggggccagca cagccctacc ccgccccagt atcatgcat ggtccccac accggttccc      1215
135 tgaacccttc ttggattaag gaagactgaa gactagcccc ttttctggga aattactttc      1275
137 ctctccctg tgtaactgg ggctgttggg gacagtgcgt gattttctcag tgatttctcta      1335
139 cagtgttgtt ccctccctca aggctgggag gagataaaca ccaaccagag aattctcaat      1395

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Input Set : A:\pto.txt  
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151 <223> OTHER INFORMATION: PCR Primer	
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159 <212> TYPE: DNA	
160 <213> ORGANISM: Artificial Sequence	
162 <220> FEATURE:	
164 <223> OTHER INFORMATION: PCR Primer	
166 <400> SEQUENCE: 5	20
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173 <213> ORGANISM: Artificial Sequence	
175 <220> FEATURE:	
177 <223> OTHER INFORMATION: PCR Probe	
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185 <212> TYPE: DNA	
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188 <220> FEATURE:	
190 <223> OTHER INFORMATION: PCR Primer	
192 <400> SEQUENCE: 7	19
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198 <212> TYPE: DNA	
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201 <220> FEATURE:	
203 <223> OTHER INFORMATION: PCR Primer	
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212 <213> ORGANISM: Artificial Sequence	
214 <220> FEATURE:	
216 <223> OTHER INFORMATION: PCR Probe	
218 <400> SEQUENCE: 9	20
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RAW SEQUENCE LISTING  
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239 <223> OTHER INFORMATION: intron 2:exon 3
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246 <222> LOCATION: (1495)...(2396)
247 <223> OTHER INFORMATION: intron 3
249 <221> NAME/KEY: exon
250 <222> LOCATION: (3213)...(3316)
251 <223> OTHER INFORMATION: exon 6
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270 cttctaataca caaatttcct tccggctgcc attttgaaag tgggccagga aatggagatg 180
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290 ggcacagagc catcttcttc aatcggatcg gtggagtga gcaggacact atcctggccg 780
292 agggccttca cttcaggtaa tggcgggcag agcctgctga ccctgacctt tcaccttga 840
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Input Set : A:\pto.txt  
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## VERIFICATION SUMMARY

PATENT APPLICATION: US/10/020,478

DATE: 01/07/2002

TIME: 20:32:08

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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
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L:237 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10  
L:241 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10  
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